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Gly Pro Ala Ser Phe Gln Cys Asn Ser Ser Thr Cys Ile Pro Gln Leu
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Trp Ala Cys Asp Asn Asp Pro Asp Cys Glu Asp Gly Ser Asp Glu Trp
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ccg cag cgc tgt agg ggt ctt tac gtg ttc caa ggg gac agt agc ccc   31269

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Pro Gln Arg Cys Arg Gly Leu Tyr Val Phe Gln Gly Asp Ser Ser Pro
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Cys Ser Ala Phe Glu Phe His Cys Leu Ser Gly Glu Cys Ile His Ser
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Ser Trp Arg Cys Asp Gly Gly Pro Asp Cys Lys Asp Lys Ser Asp Glu
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Thr Leu Asp Lys Val Cys Asn Met Ala Arg Asp Cys Arg Asp Trp Ser
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Thr Glu Val Ala Ser Asn Arg Ile Tyr Trp Ser Asp Leu Ser Gln Arg
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atg atc tgc ag gtgagcgtcg cccctgcctg cagccttggc ccgcaggtga   38594
Met Ile Cys se
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gatgagggct cctggcgctg atgcccttct ctcctcctgc ctcag c acc cag ctt   38649
                                   r Thr Gln Leu
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gac aga gcc cac ggc gtc tct tcc tat gac acc gtc atc agc aga gac   38697
Asp Arg Ala His Gly Val Ser Ser Tyr Asp Thr Val Ile Ser Arg Asp
        440   445   450

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atc cag gcc ccc gac ggg ctg gct gtg gac tgg atc cac agc aac atc   38745
Ile Gln Ala Pro Asp Gly Leu Ala Val Asp Trp Ile His Ser Asn Ile
        455   460   465

tac tgg acc gac tct gtc ctg ggc act gtc tct gtt gcg gat acc aag   38793
Tyr Trp Thr Asp Ser Val Leu Gly Thr Val Ser Val Ala Asp Thr Lys
      470   475   480

ggc gtg aag agg aaa acg tta ttc agg gag aac ggc tcc aag cca agg   38841
Gly Val Lys Arg Lys Thr Leu Phe Arg Glu Asn Gly Ser Lys Pro Arg
      485   490   495

gcc atc gtg gtg gat cct gtt cat gg gtgcgtatcc acgacgctga 38887
Ala Ile Val Val Asp Pro Val His gl
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